

FIG. 1A

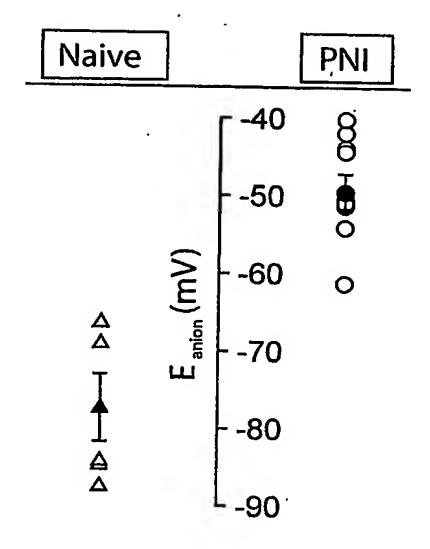


FIG. 1B

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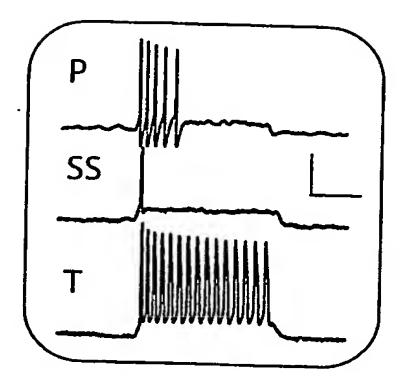


FIG. 1C

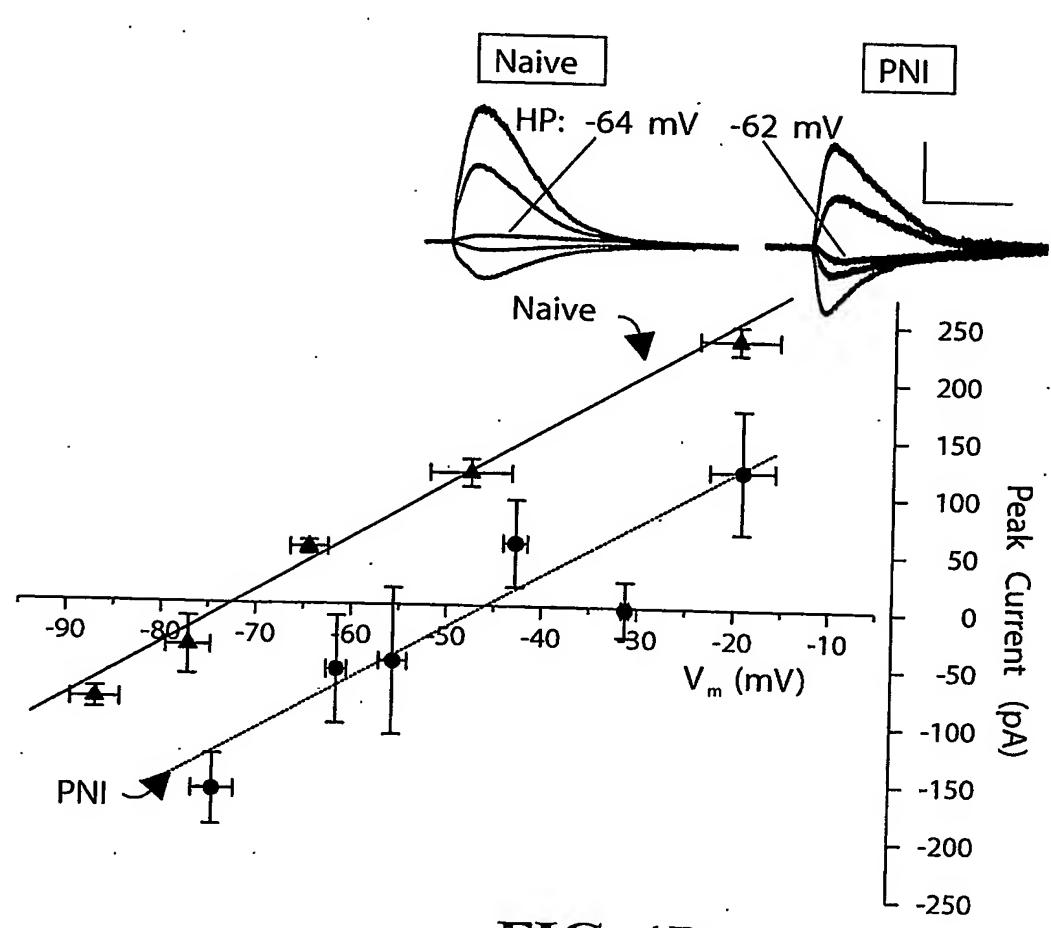
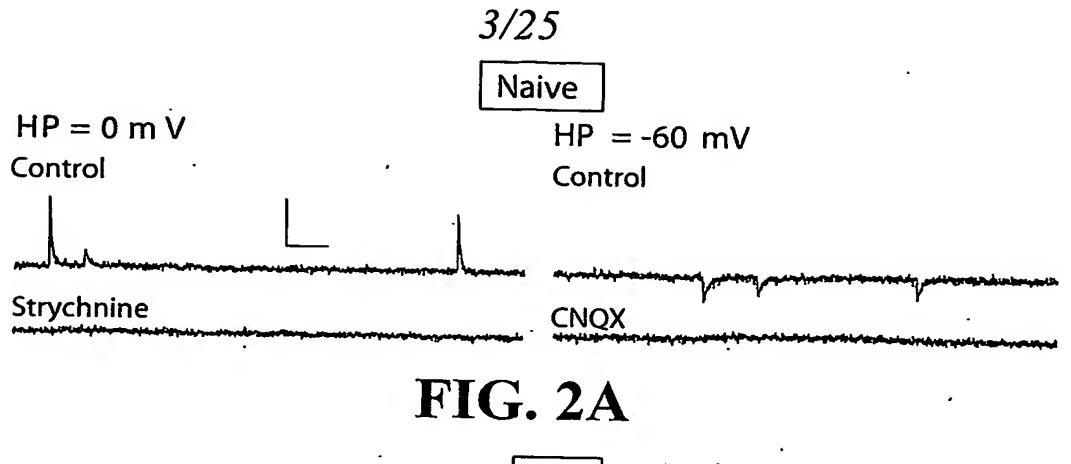


FIG. 1D



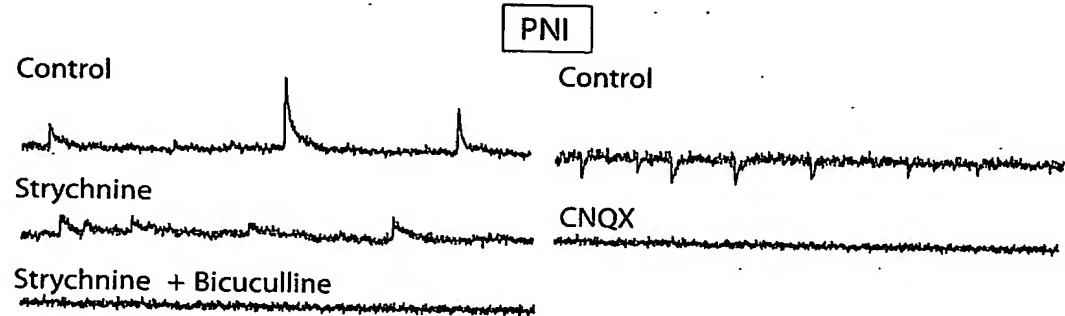


FIG. 2B

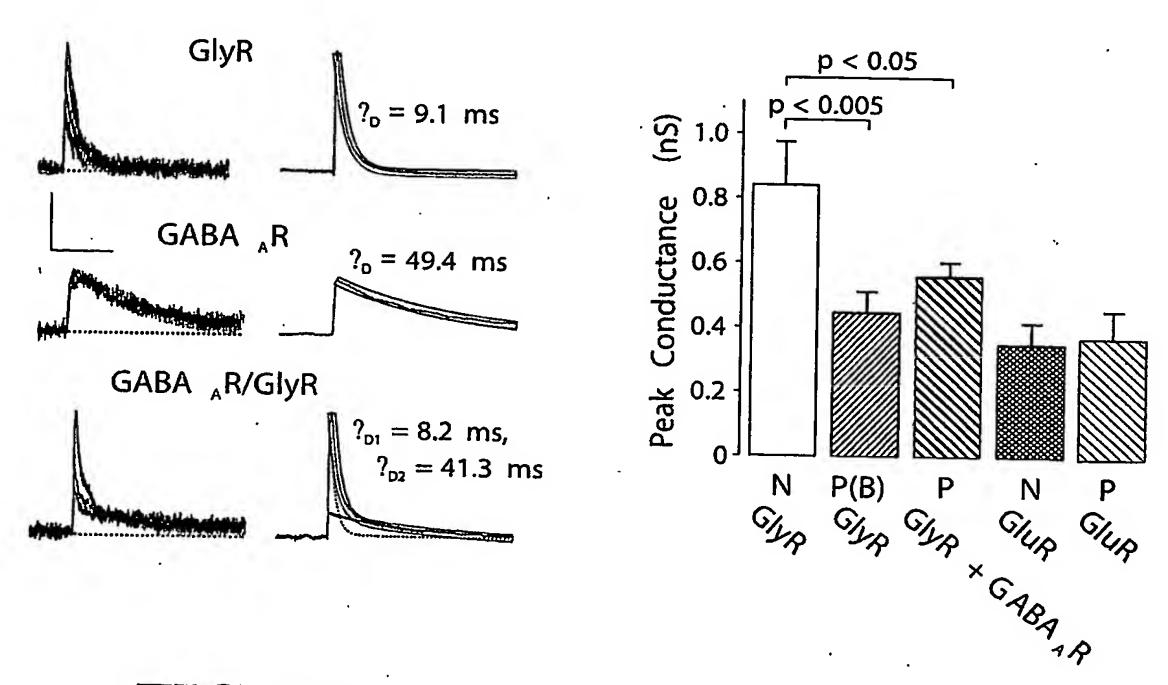


FIG. 2C

FIG. 2D



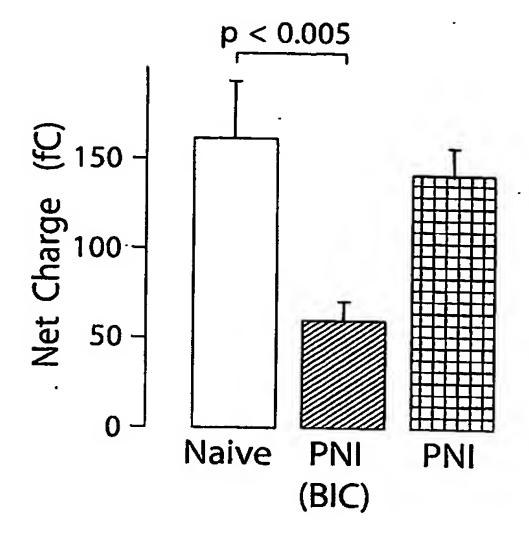


FIG. 2E

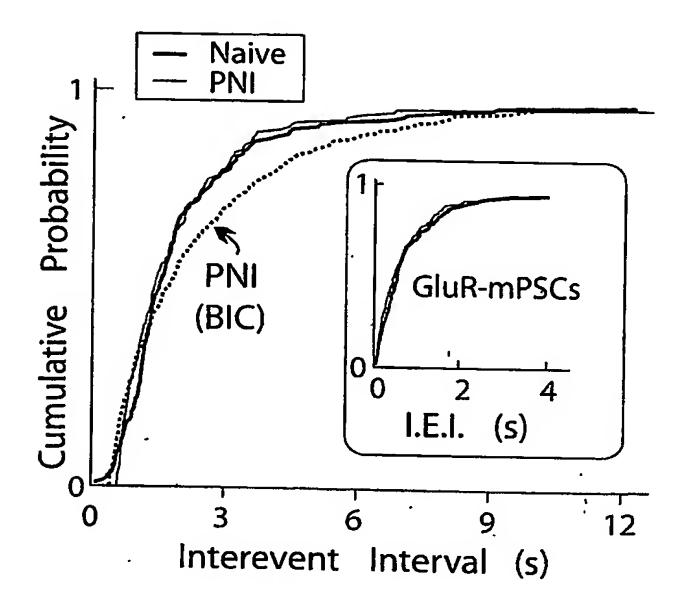


FIG. 2F



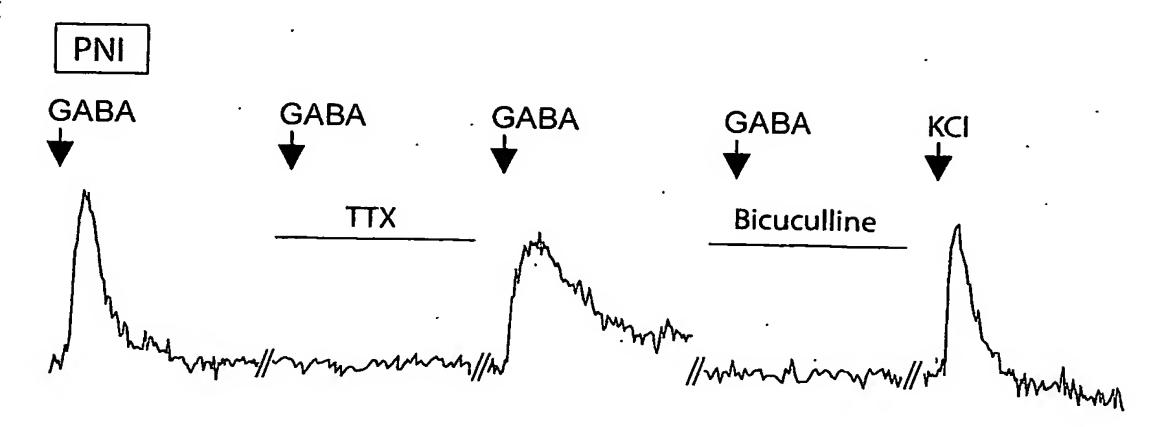


FIG. 3A

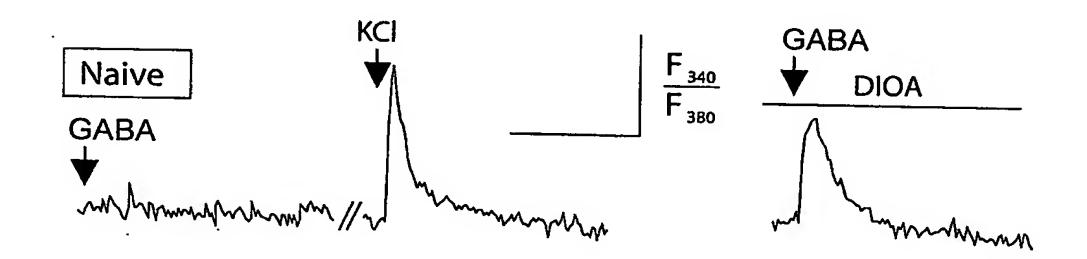


FIG. 3B

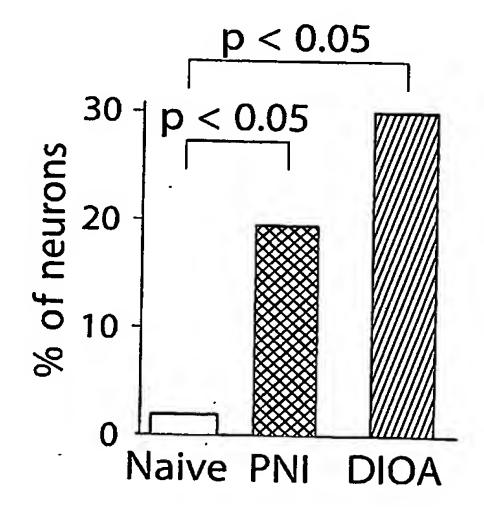
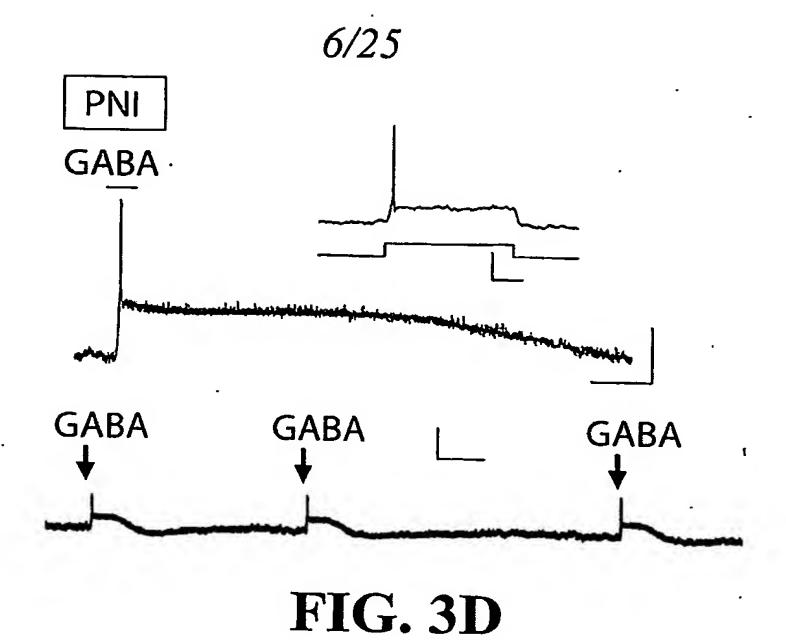


FIG. 3C



Suprathreshold

* Subthreshold

* Bicuculline

FIG. 3E

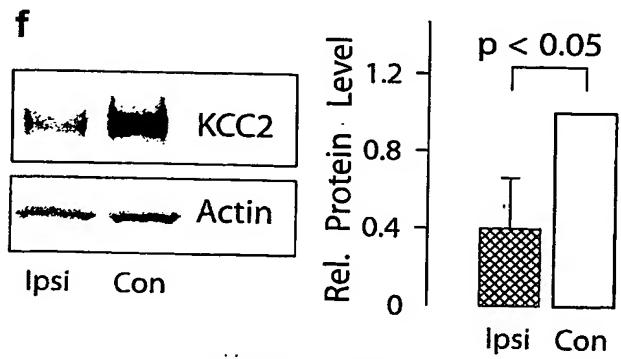
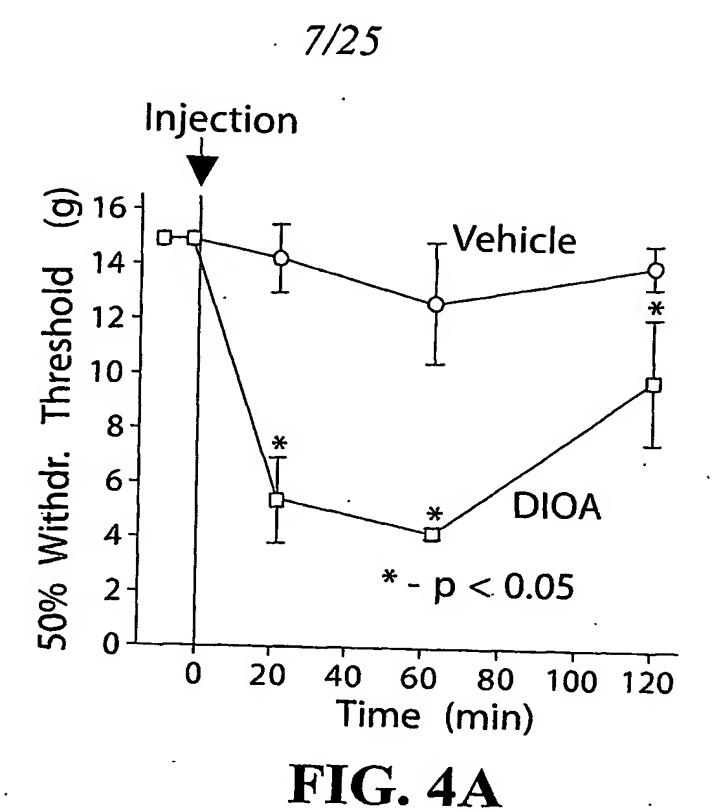


FIG. 3F



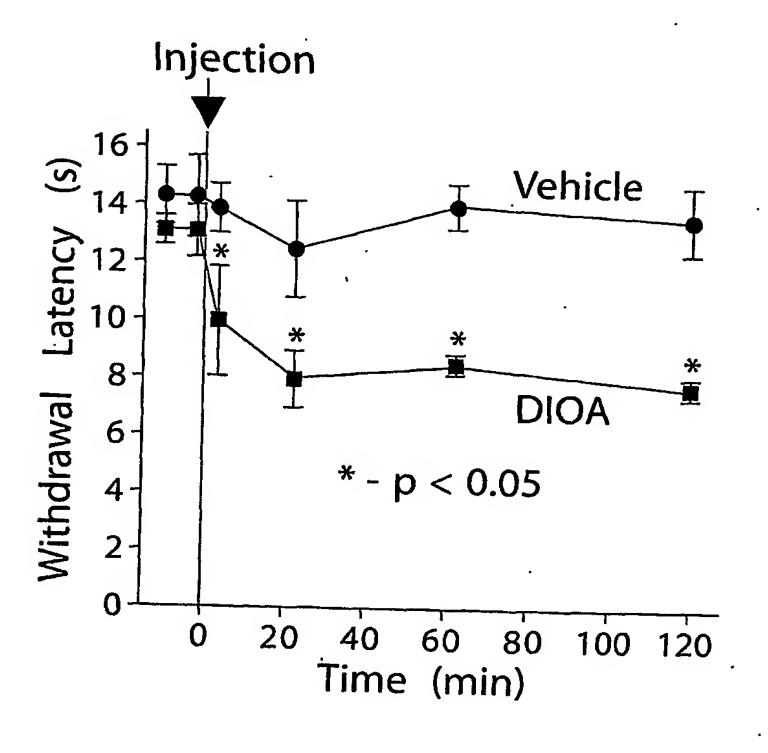


FIG. 4B



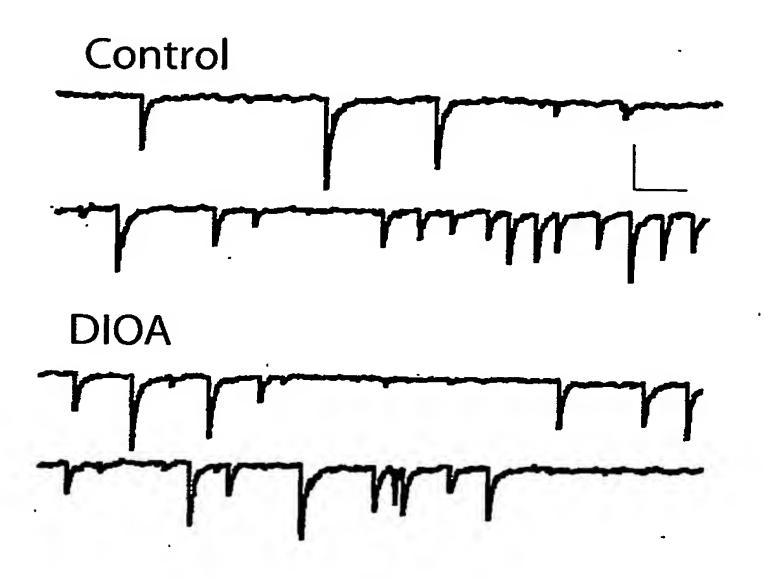


FIG. 4C

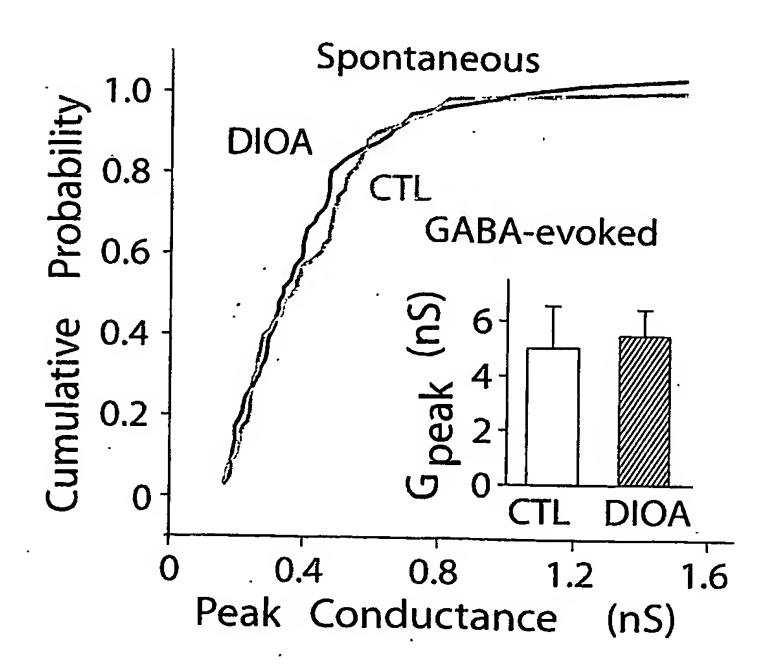


FIG. 4D

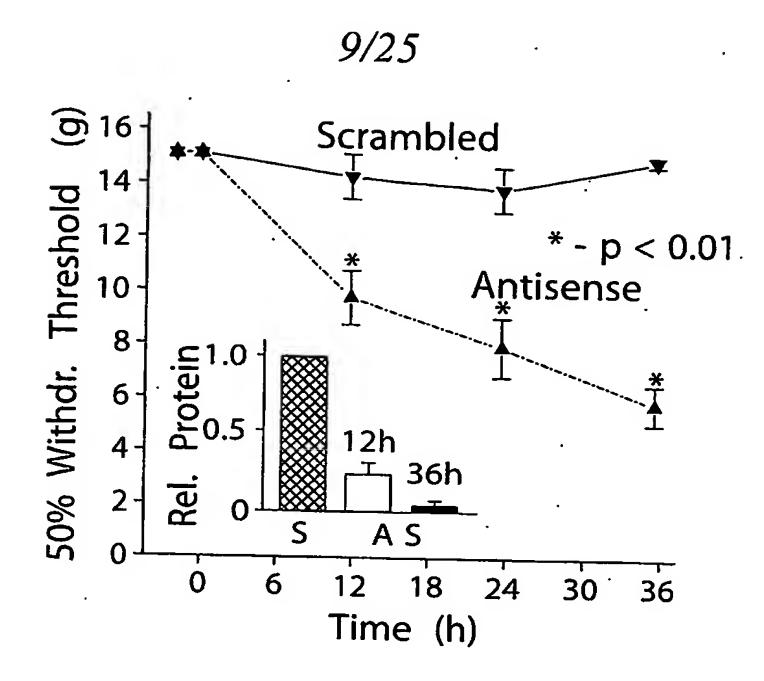


FIG. 4E

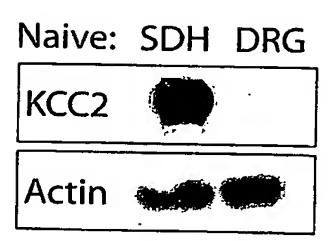


FIG. 4F

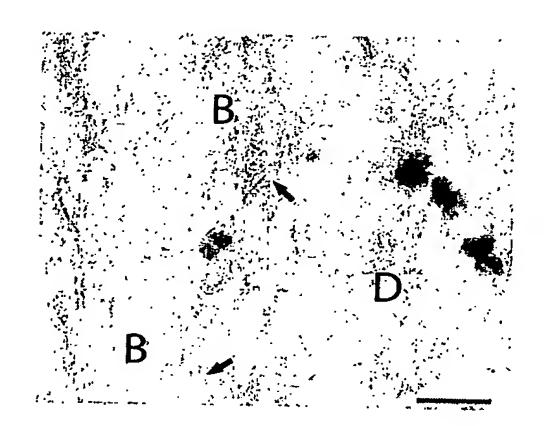


FIG. 4G

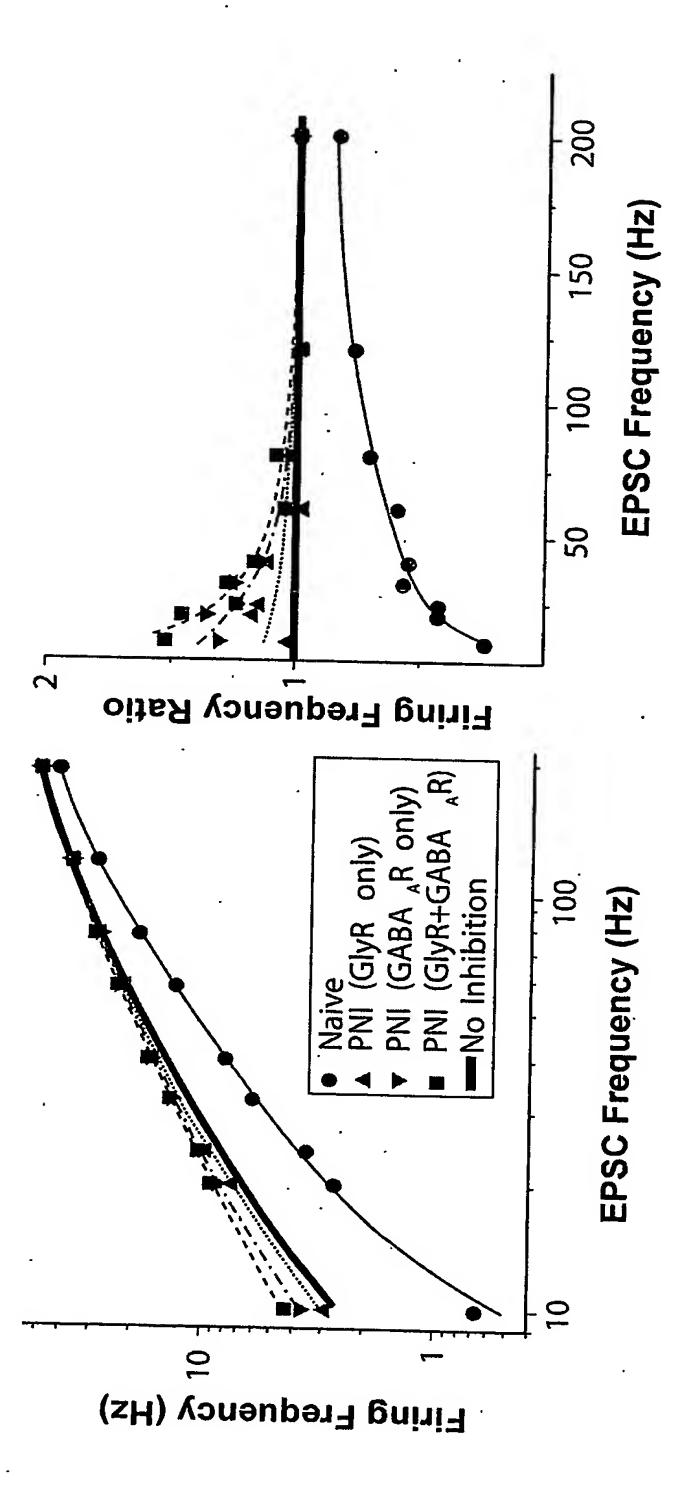


FIG. 5A

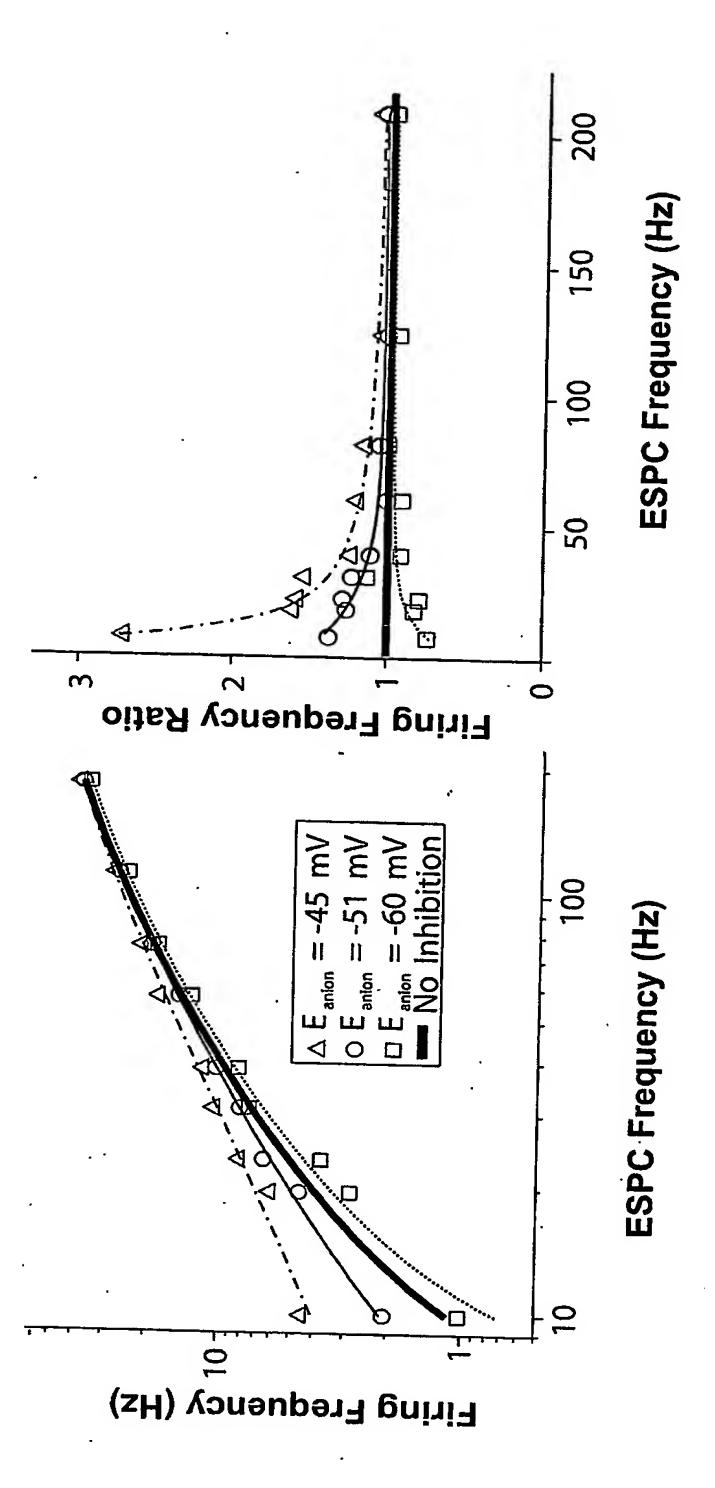


FIG. 5B

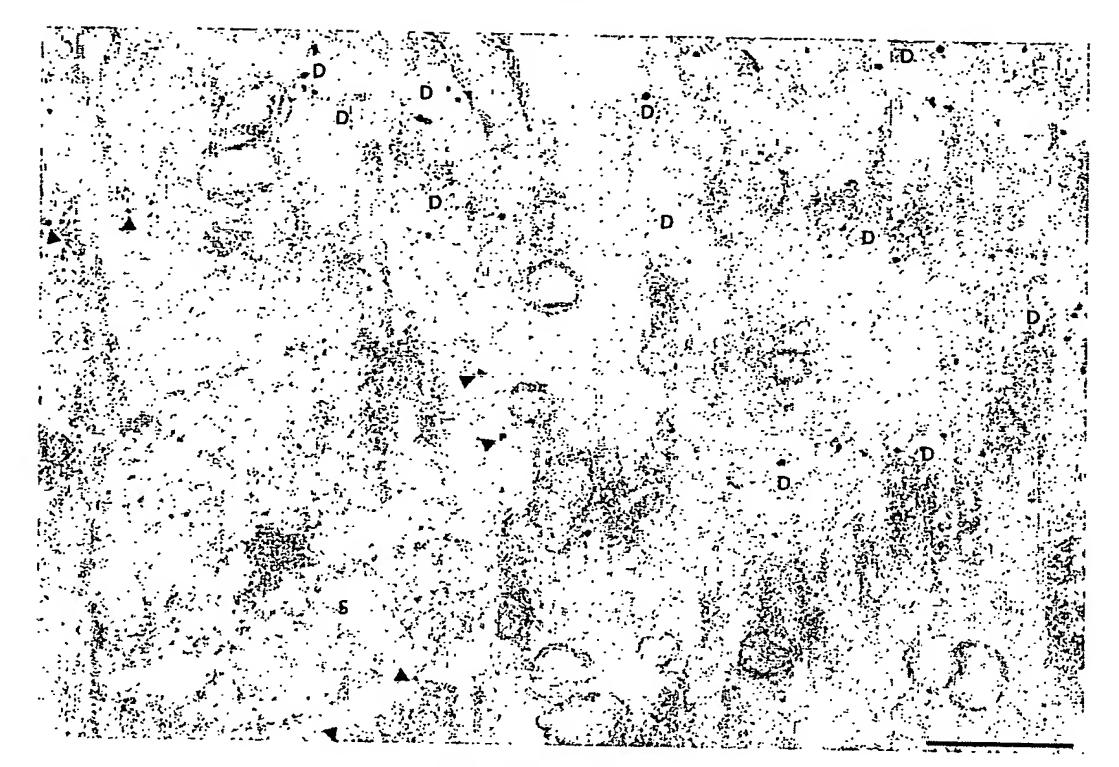


FIG. 6A

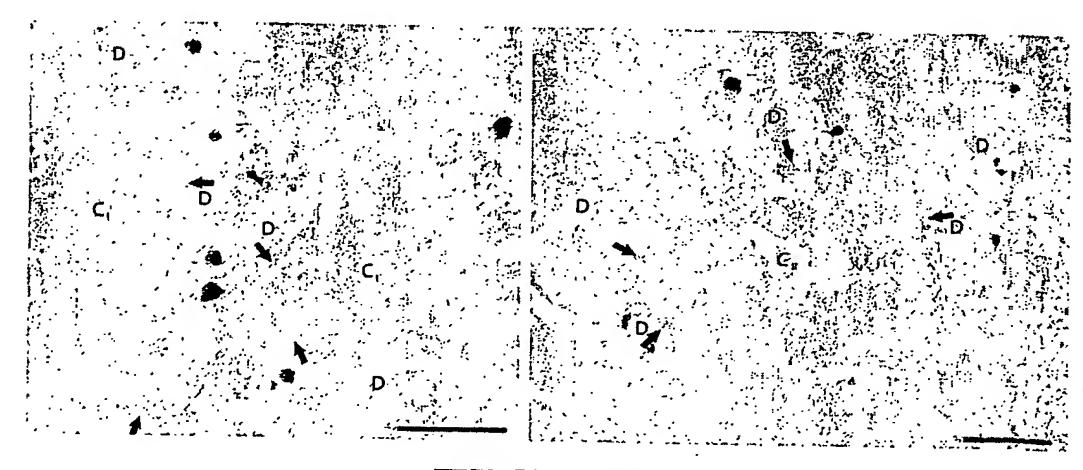


FIG. 6B

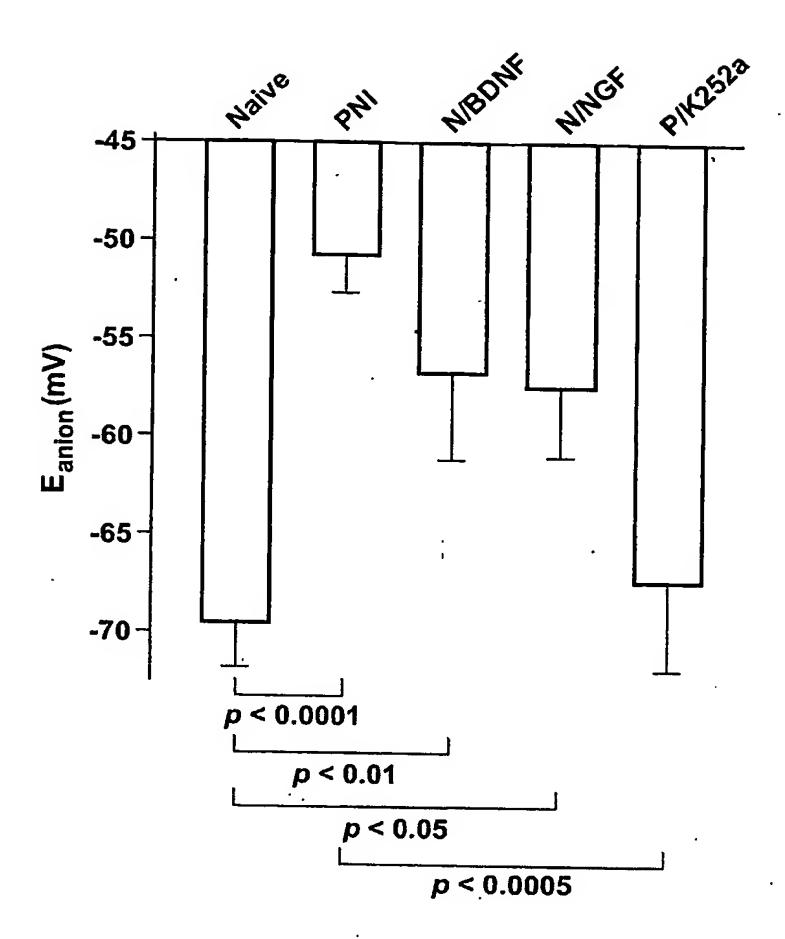


FIG. 7

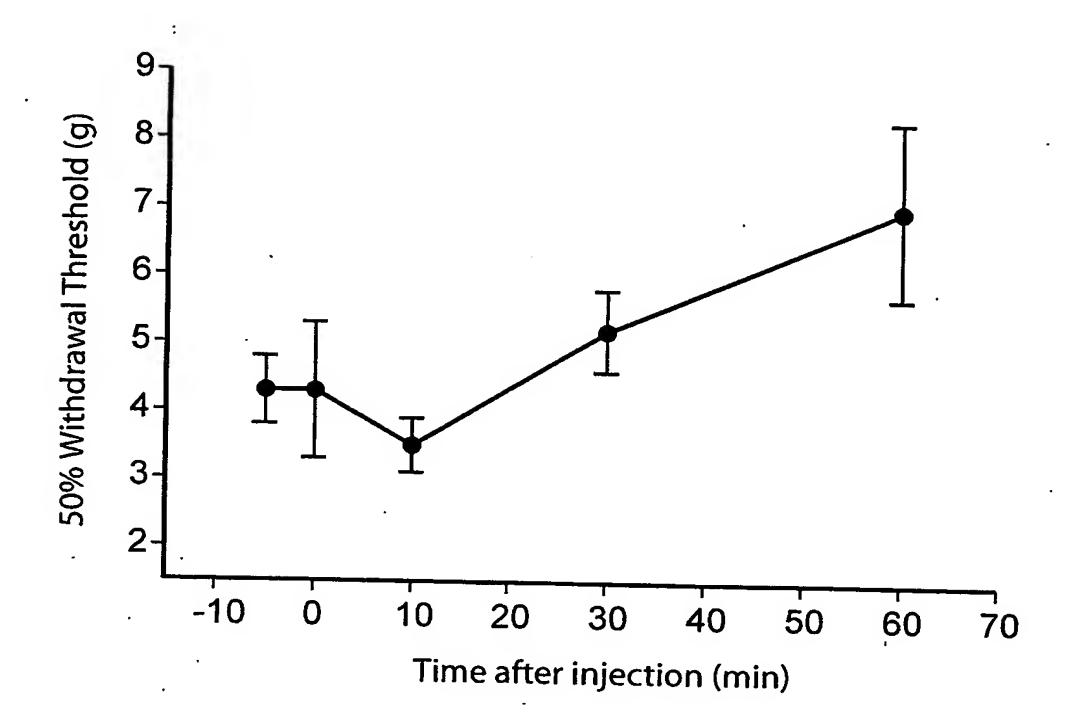


FIG. 8

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Human KCC2 polypeptide and DNA sequences

(Mount, D.B. and Song, L. (2002) Brain Res. Mol. Brain Res. 103 (1-2), 91-105; ACCESSION: AF208159)

Human KCC2 polypeptide (SEQ ID NO:2):

MPNNLTDCEDGDGGANPGDGNPKESSPFINSTDTEKGKEYDGKN

MALFEEEMDTSPMVSSLLSGLANYTNLPQGSREHEEAENNEGGKKKPVQAPRMGTFMG VYLPCLQNIFGVILFLRLTWVVGIAGIMESFCMVFICCSCTMLTAISMSAIATNGVVP AGGSYYMISRSLGPEFGGAVGLCFYLGTTFAGAMYILGTIEILLAYLFPAMAIFKAED ASGEAAAMLNNMRVYGTCVLTCMATVVFVGVKYVNKFALVFLGCVILSILAIYAGVIK SAFDPPNFPICLLGNRTLSRHGFDVCAKLAWEGNETVTTRLWGLFCSSRFLNATCDEY FTRNNVTEIQGIPGAASGLIKENLWSSYLTKGVIVERSGMTSVGLADGTPIDMDHPYV FSDMTSYFTLLVGIYFPSVTGIMAGSNRSGDLRDAQKSIPTGTILAIATTSAVYISSV. VLFGACIEGVVLRDKFGEAVNGNLVVGTLAWPSPWVIVIGSFFSTCGAGLQSLTGAPR LLQAISRDGIVPFLQVFGHGKANGEPTWALLLTACICEIGILIASLDEVAPILSMFFL MCYMFVNLACAVQTLLRTPNWRPRFRYYHWTLSFLGMSLCLALMFICSWYYALVAMLI AGLIYKYIEYRGAEKEWGDGIRGLSLSAARYALLRLEEGPPHTKNWRPQLLVLVRVDQ DQNVVHPQLLSLTSQLKAGKGLTIVGSVLEGTFLENHPQAQRAEESIRRLMEAEKVKG FCQVVISSNLRDGVSHLIQSGGLGGLQHNTVLVGWPRNWRQKEDHQTWRNFIELVRET TAGHLALLVTKNVSMFPGNPERFSEGSIDVWWIVHDGGMLMLLPFLLRHHKVWRKCKM RIFTVAQMDDNSIQMKKDLTTFLYHLRITAEVEVVEMHESDISAYTYEKTLVMEQRŞQ ILKQMHLTKNEREREIQSITDESRGSIRRKNPANTRLRLNVPEETAGDSEEKPEEEVQ LIHDQSAPSCPSSSPSPGEEPEGEGETDPEKVHLTWTKDKSVAEKNKGPSPVSSEGIK DFFSMKPEWENLNQSNVRRMHTAVRLNEVIVKKSRDAKLVLLNMPGPPRNRNGDENYM **EFLEVLTEHLDRVMLVRGGGREVITIYS**

Human KCC2 DNA (SEQ ID NO:1):

1 atgcccaaca acctgacgga ctgcgaggac ggcgatgggg gagccaaccc gggtgatggc 61 aaccccaagg aaagcagtcc cttcatcaac agcaccgaca cagagaaggg aaaggagtat 121 gatggcaaga acatggcctt gtttgaggag gagatggaca ccagccctat ggtgtcctcc 181 ttgctcagtg gcctggccaa ctacaccaac ctgcccagg gaagtaggga gcatgaagag 181 gcagaaaaca atgagggtgg aaaaaaagaag ccggtgcagg ccccacgcat gggcaccttc 181 acctgggtgg acctgccgtg cctgcagaac atctttggcg tcatcctct cctgcggctc acctgcgtgg tgggcattgc aggcatcatg gagtccttct gcatggtgt catctgctgc

421	tcctqtacqa	tactcacaac	catctccatc	acteonatt.		
481	gctggtggct	cctactacat	gatttccacg	agtgcaattg	caacgaatgg	tgttgtgcct
541	ggcctctqct	tctacctor	cactacettt	general and a	cagagtttgg	gggtgccgtg gggcaccatc
601	gaaatcctgc	tagettaget	cttcccacc	atggccatct	tgtacatcct	gggcaccatc
661	ggggaggcag	cagccatact	gaacaacato	acygedatet	tcaaggcaga	agatgccagt
721	atggccactg	tagtatttat	gaacaacacg	cgtgtttacg	gcacctgtgt	gctcacctgc
781	gattatatca	tecteteest	cctcccate	tatgtcaaca	agtttgccct	tgtcttcctg
841	ccacccaact	tecegateta	cctagtage	tatgctgggg	tcatcaagtc	tgccttcgac
901	gtctgtgcca	agetggettg	ggaagaaat	aaccgcacgc	tgtctcgcca	tggctttgat
961	ttctgctcct	ctcacttcct	ggaaggaaac	gagacggtga	ccacacggct	atggggcctt
1021	acagagatics	agggettee	tagtagtage	tgtgatgaat	acttcacccg	aaacaatgtc
1081	tectacetga	ccaaggggggg	cattatasa	agtggcctca	tcaaagagaa	cctctggagc
1141	gatggcactc	ctategagge	gactgtggag	aggagtggga	tgacctcggt	gggcctggcc
1201	accetactor	ttagcatcta	gyaccaccet	tatgtcttca	gtgatatgac	ctcctacttc
1261	tctggggacc	taggaataa	ggagaactea	gtcacaggga	tcatggctgg	ttctaaccgc
	5555566	CHARRENES	CCayaadica	ATCCCCAC+~	~~~~~~~~~	
		o cyclatat	Cayclecare	GTTCTATTTA	acacache an b	A
	2	ucaageeegg	CHARACTOTO	aarggcaacc	+ 00+00+000	
	-33000000	- vwcgggcaat	Lucalcuda		aaaaatataa	and the second s
	3-5-0-0-4		acuccuacta	Caddecatet	0020001taa	and the second of the second o
		- Letter Laggera	LYYCAAGGCC	aatooadaa	AASAAt AAA	
		- ccgcgagac	Lydualcete	attocatece	tagaaaaat	*****
		CCCCCCCCCCC	utuctacard	TTTMTMaaka	+~~~~+	
	33-5	- Jacobaabby	4444CCaccc	LLLCGAtatt	ancactors	والمراجع بالمراجع والمراجع
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	3-04030004	GC - GG GC C	catctacaac	じみぐみじじべっぺし	3000teen	
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	773		- caadaaci.uu	ACCCCACACA	+~~+~~+ ~~+	
	3	ugaatgtqqt	ucaccccaa	CECCECECS	+~~~~~~~	- · L ·
_		- Jacourty	999666666	CEEGAGGGGA	へのたたたのだみべっ	
		コングしいりゅんけん	4LULALUAGO	COCCEGATAA	200020000	
	2222223	- Jac-	CLCCAACEEG	COTOSTOOO	+~+~~~~+~ ~	man da la
	222220000	BBBBBCCBCa	ycacaacact	aracttatta	aataaaaaa	
*		accaccagac	чьччаччаас	CECATTOROS	taataaaaa	*****
			Caccaadaac	GTTTCCatct	ttaataaaa	
	77777777	geageacega	Cycliagia	ALLGLGGGAGG	atararani	
		~300gcgca	CCacaauulc	LUGCGGAAAt	777277+~~~	
. – –	2-33-4-423	- Julyacaa	Layualcead	atgaagaagg	atotoacoa	
_		- June Lycyda	MACCAGGG	araaaaraa	21/22/24/24	
		uguugucgee	ggugalqqaq	cadediteee	adatectes	202021
		~~5~5~554	Seggadate	cadadratca	cacatoacto	200200-b-
	acceggagaa	agaacccage	caacacqcqq	Cliccacataa	acateceans	2020250
	334344369	uugagaagcc	ayayyaddad	atacaaataa	teesees	man mela au auta au au au
	-35C-C	grageteete	ulccccadaa	gaggagggta	200000000	
		Guulla	CLUUACCAAA	7272277 AAA	+~~~~~~~	_ 1
		TOUCELINE	MAMCALCAAG	OBCETCTECS.	MMStanaaa.	
		-Jedouacyc	MCMACACALA	cacacoooco	TMMMMM ANN	
	2-2-2-4		Caagettget	LEGCECAACA	tacatacaca	togggggggg
	-353-59	~cyaaaacta	Catudadete	CECGAGGERR	tagagagaga	
	5-5-555		Lygucuadad	gtcatcacca	totactooto	20222000
		コココーししらははい	acaccaacc	CGCGGCTCCC	asaaaat aaa	~~~~~~~
	3-03-03-04	Cogocoacac	acadaceeta	Edcccatata	atavaaaaht	
	3	JUDDOU EENE	CCALLAGGGC	TOSTTONASA.	200000000	
	J J 5 - C	occus cycc	auttluaccc	CEGGGTC++	aataaaatt.	
	22-02000		gacyclycaa	taaaggrege	CACCACACA	~~~~~
	77~7~73333		CCCCaddlad	TCCATGCGGG	agattagtag.	والمستحدث والمستحدث
		coccogccc	ACACTCCECC	aacactacta	aataaataaa	~~~~~~
·		PEDDEEEC	CCACACCEAC	COLCLCCCCC	aaaaattata	
つりみ丁	aaaacacaaa	cggccgagcc	tatacatagt	gtacaggaga	catcgcgtgt	atttttaaca
			•			- 3

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3901 tececatatt tatgtgaeta gaagegeaae agaetteteg ceatagtega geteteege
3961 tggggggact gcgggggggc gaggcctcgg gaagctgaat tttccttgac gtccaagagt
4021 ttgagagcga aagtgcttta ggcccaggcg ggggtcgtgg cctcgttccc tcgacacctc
4081 cgtcctgctc tcgcctcttc gccctttccg cgcgcccttg gcttcccacc ctcctcca
4141 gtccttttcc gagatgaggt gagacaaggg tccaactttt cctggattcg cctcccagcg
4201 gacgtgaget tecaetgegg etgeagagae gegageaace tetteteate ggetettatg
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4321 gccccaccga ggaagccccg ccccggtgcc ttcgctgggg agcaggcgtc tctcctcagt
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4441 tggagggcgc cccctccccg gagtttcctc cctgggacaa gtgagggagg aggggccga
4501 ttctggttta ggggccggac ccactgagag gccccagagc cgcccgtgat gttcctccc
4561 cgtccccatc tggcagctcc tgtctcgcct gagggaccca gccgccttct ccgtgctctg
4621 gggccgggcc tcgctgctta gcagcggcct ctagctccgt ctcccgggga cctgggcctg
4681 agggaggget ggagteagea egegetttgt eettagegee tgtetgetet eetetaaeta
4741 ggacccaggg cetttggett ceccagetea teettggeeë tteegeteea ceageetggt
4801 ctgaggcgtg ctctgtcctt agagaaggcg cggtggccgg gttcccttcc cctagggcac
4861 attactaagg gggtcaggca ctgcatgctc gttccagcac catctgggac tgggtacagt
4921 acctccagcc ccagggccct gacctgcgca cctagcttga catctcacgc acctcccaga
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5041 ctagagctgg agctggcgcc acccagacag cgtcaggtgt ggctggggta ggtttggagg
5101 tctgccagtt acgccaagtc ccctctgaga ttcgatcagg ggactggata gattctttca
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5341 tctgggccca gattgtctgg ttggcaagag caaagtttcc gttgatgaaa cagacatccc
5401 acaacaaaaa cccaagtttt ctgtgctaca tgtgcaatat ttgttatgaa tgttatcaca
5461 agtcattcat caagttatct ttataatcac tgtagttaga tgtttcatgt ccattcaagt
5521 gacttttatt ctgagtgcaa tatttcaata gccttgtagt gataactagt gttgcttttg.
5581 tttagatgat ctatgtgcag ggcaatgcaa tgaagttgaa accccttggt aataggagag
5641 gttgcaaacc aaatcaagag tatttattac tattactgct attattatta ggcctgcctt
5701 taattttcag tgtaagtgtt cagtatgccg catcctgcct cagtattgat cttgtgttct
5761 ttgtgccaat atgaaaagga gagggttggt tctttccttt attgttgaat gctcccattt
5821 aatgetttat ggettttact gtattacttt tttagaetee egtetgeaca aaatgeaata
5881 aaaataattt tattataaaa aaaaaaa
```

FIG. 9 (Continued)

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Mouse KCC2 (K-Cl cotransporter [Slc12a5]) polypeptide and DNA sequences

(Ehringer, M.A., et al. (2001) Mamm. Genome 12 (8), 657-663; ACCESSION: AF332064)

Mouse KCC2 polypeptide (SEQ ID NO:4):

MLNNLTDCEDGDGGANPGDGNPKESSPFINSTDTEKGREYDGRN

MALFEEEMDTSPMVSSLLSGLANYTNLPQGSREHEEAENNEGGKKKPVQAPRMGTFMG VYLPCLQNIFGVILFLRLTWVVGIAGIMESFCMVFICCSCTMLTAISMSAIATNGVVP AGGSYYMISRSLGPEFGGAVGLCFYLGTTFAGAMYILGTIEILLAYLFPAMAIFKAED ASGEAAAMLNNMRVYGTCVLTCMATVVFVGVĶYVNKFALVFLGCVILSILAIYAGVIK SAFDPPNFPICLLGNRTLSRHGFDVCAKLAWEGNETVTTRLWGLFCSSRLLNATCDEY FTRNNVTEIQGIPGAASGLIKENLWSSYLTKGVIVERRGMPSVGLADGTPVDMDHPYV FSDMTSYFTLLVGIYFPSVTGIMAGSNRSGDLRDAQKSIPTGTILAIATTSAVYISSV VLFGACIEGVVLRDKFGEAVNGNLVVGTLAWPSPWVIVIGSFFSTCGAGLQSLTGAPR LLQAISRDGIVPFLQVFGHGKANGEPTWALLLTACICEIGILIASLDEVAPILSMFFL MCYMFVNLACAVQTLLRTPNWRPRFRYYHWTLSFLGMSLCLALMFICSWYYALVAMLI AGLIYKYIEYRGAEKEWGDGIRGLSLSAARYALLRLEEGPPHTKNWRPQLLVLVRVDQ DQNVVHPQLLSLTSQLKAGKGLTIVGSVLEGTFLDNHPQAQRAEESIRRLMEAEKVKG FCQVVISSNLRDGVSHLIQSGGLGGLQHNTVLVGWPRNWRQKEDHQTWRNFIELVRET TAGHLALLVTKNVSMFPGNPERFSEGSIDVWWIVHDGGMLMLLPFLLRHHKVWRKCKM RIFTVAQMDDNSIQMKKDLTTFLYHLRITAEVEVVEMHESDISAYTYEKTLVMEQRSQ ILKQMHLTKNEREREIQSITDESRGSIRRKNPANPRLRLNVPEETACDNEEKPEEEVQ LIHDQSAPSCPSSSPSPGEEPEGERETDPEVHLTWTKDKSVAEKNKGPSPVSSEGIKD FFSMKPEWENLNQSNVRRMHTAVRLNEVIVNKSRDAKLVLLNMPGPPRNRNGDENYME FLEVLTEQLDRVMLVRGGGREVITIYS

Mouse KCC2 DNA (SEQ ID NO:3):

1 gagcaagcga gcgagcggag aaggcgggca gaggggcgcg ggcgaagcgg cgcagccatc 61 ccgagcccgg cgccgcgag ccaccatgct caacaacctg acggactgcg aggacggcga 121 tgggggagcc aaccccggtg atggcaaccc caaagagagc agtcccttca tcaacagcac 181 ggacaccggag aagggcagag agtacgatgg caggaacatg gccctgtttg aggaggagat 241 ggacaccagc cccatggtat cctccctgct cagtgggctg gccaactaca ccaacctacc 301 ccagggaagt agagagcatg aagaagcaga aaataatgag ggtggaaaaa agaagccggt

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361 gcaggctcct cgaatgggca ccttcatggg.tgtgtacctg ccgtgcctgc agaacatctt
  421 tggtgtcatc ctcttcctgc ggctcacgtg ggtggtgggc atcgcgggca tcatggagtc
  481 cttctgtatg gtcttcattt gctgctcctg tacgatgctc acagccattt ccatgagtgc
  541 aatcgcaacc aatggtgttg tgcctgctgg tggctcgtac tacatgattt ccaggtctct
  601 gggcccggag tttggggcg ccgtgggcct ctgcttctac ctgggcacca cctttgctgg
  661 ggctatgtac atccttggca cgatcgagat cctgctggct tatctcttcc cagctatggc
  721 catcttcaag gcagaagatg ccagtgggga ggcggccgcc atgctgaaca acatgcgggt
  781 gtatggcacc tgtgtgctca cctgcatggc caccgttgtc tttgtgggtg tcaagtacgt.
  841 caacaagttt gccttggtct tcctgggttg egtcatcctg tccatcctgg ccatctatgc
  901 aggggtcatc aagtctgcct tcgacccacc caatttcccg atctgcctcc tggggaaccg
  961 cacgctgtct cgccatggct ttgatgtctg tgccaagctg gcttgggaag gaaatgagac
 1021 agtgaccaca cggctctggg gccttttctg ctcctcccgc ctcctcaatg ccacctgtga
 1081 tgagtacttc acccgaaaca atgtcacaga gatccagggc attcctggtg ctgccagtgg
 1141 tctcatcaaa gagaacctgt ggagttctta cctgaccaaa ggggtgattg tcgagaggcg
 1201 tgggatgccc tctgtgggcc tggcagacgg tacccccgta gacatggacc acccctatgt
 1261 cttcagtgat atgacctcct acttcaccct gctcgttggt atctacttcc cctcagtcac
1321 agggatcatg gctggctcaa accgatctgg agacctgcgg gatgcccaga agtctatccc
1381 tactggaact atcctggcca ttgctaccac ctctgctgtc tacatcagct ctgttgttct
1441 gtttggagcc tgcatcgagg gggtcgtctt acgggacaag tttggggaag ctgtgaatgg
 1501 caacttggtg gtgggcaccc tggcctggcc ttctccctgg gtcatcgtca taggctcttt
1561 cttctctacc tgtggggctg gattacagag cctcacaggg gccccacgtc tgctgcaggc
1621 catctcccgg gatggcatag tgcccttcct gcaggtcttt ggccatggca aagctaatgg
1681 agagccaacc tgggcgctgc tgctgactgc ctgcatctgt gagatcggca tcctcatagc
1741 ctccctggat gaggtcgccc ctatactttc catgttcttc ctaatgtgtt acatgtttgt
1801 gaacttggct tgtgcggtgc agacgctgct gaggacaccc aactggaggc cacgatttcg
1861 ctattaccac tggactctct ccttcctggg catgagcctc tgcctggccc tcatgttcat
1921 ttgctcctgg tactacgcac tggtggccat gctcattgcc ggactcattt ataagtacat
1981 cgagtaccgg ggggcggaga aggagtgggg ggatggaatc cgaggcctgt ctctcagtgc
2041 agcacgctat gctctcttgc gcctggagga aggacctccg catacgaaga actggaggcc
2101 ccagctgctg gtgctggtgc gtgtggacca ggatcagaac gtggtgcatc cgcagctgct
2161 ctccctgacc tcccagctca aggcagggaa gggcctgacc attgtgggct ccgtccttga
2221 gggcaccttt ctggacaacc atccacaggc tcagcgggca gaggagtcta tcaggcgcct
2281 gatggagget gagaaggtga agggettetg ceaggtagtg atetecteea acetgegtga
. 2341 tggtgtgtcc cacctgatcc agtctggggg cctcggggga ttgcaacaca ataccgtgct
2401 ggtgggctgg cctcgcaact ggaggcagaa ggaggatcat cagacatgga ggaacttcat
2461 cgaactggtc cgggaaacta cagccggcca cctcgccctg ctggtcacca agaatgtttc
2521 catgtttccc gggaaccctg agcgcttctc ggagggcagc attgacgtgt ggtggattgt
2581 gcacgacggg ggcatgctca tgctgctgcc cttcctgctg cgacaccaca aggtctggag
2641 gaaatgcaaa atgcggatct tcaccgtggc ccagatggac gataacagta tccagatgaa
2701 gaaggacctg accacgtttc tgtaccactt acgcattact gcagaggtgg aggtggtaga
2761 gatgcatgag agcgacatct cggcatacac ctacgagaag acattagtaa tggagcaacg
2821 atctcagatc ctcaaacaga tgcacctcac caagaacgag cgggaacggg agatccagag
.2881 catcacagac gagtctcggg gctccattcg gaggaagaat ccagccaacc cccggctccg
2941 cctcaatgtt cccgaagaga cagcgtgtga caatgaggag aagccagagg aggaggtgca
3001 gctgatccat gaccagagtg ctcccagctg ccctagcagc tcgccatctc caggggagga
3061 gcccgagggg gagagggaga cagacccaga ggtgcatctt acctggacca aggataagtc
3121 agtggcagag aagaataaag gccccagtcc cgtctcctcc gagggcatca aggacttctt
3181 cagcatgaag ccggagtggg aaaacttgaa ccagtccaat gtacggcgca tgcacacagc
3241 tgtgcggctg aacgaggtca tcgtgaataa atctcgggat gccaagctag ttttgctcaa
3301 catgcccggg cctccccgca accgcaatgg ggatgaaaac tacatggaat tcttggaggt
3361 cctcactgag caactggacc gggtgatgct ggtccgcggt ggcggccgag aggtcatcac
3421 catctactcc tgaaggccag gacctgccac tccggcccga gcgcgcccgg cccgcggccc
3481 ccagagccct cgccgccct ccccgccgct gtcaccgttt acataagacc cagttgccca
3541 tgccctggcc cctttccttc ccgctgcctg cagccctgag gccttgcccg tcggggctga
  3601 cccgcagggc ggcccgtgag gccccttttc tgagcctggc ctcgcccgc cggagc
```

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Rat KCC2 polypeptide and DNA sequences

(Payne, J.A., et al., (1996) J. Biol. Chem. 271 (27), 16245-16252; Gillen, C.M., et al., (1996) J. Biol. Chem. 271 (27), 16237-16244; ACCESSION: U55816)

Rat KCC2 polypeptide (SEQ ID NO:6):

MLNNLTDCEDGDGGANPGDGNPKESSPFINSTDTEKGREYDGRN

MALFEEEMDTSPMVSSLLSGLANYTNLPQGSKEHEEAENNEGGKKKPVQAPRMGTFMG VYLPCLQNIFGVILFLRLTWVVGIAGIMESFCMVFICCSCTMLTAISMSAIATNGVVP AGGSYYMISRSLGPEFGGAVGLCFYLGTTFAGAMYILGTIEILLAYLFPAMAIFKAED ASGEAAAMLNNMRVYGTCVLTCMATVVFVGVKYVNKFALVFLGCVILSILAIYAGVIK SAFDPPNFPICLLGNRTLSRHGFDVCAKLAWEGNETVTTRLWGLFCSSRLLNATCDEY FTRNNVTEIQGIPGAASGLIKENLWSSYLTKGVIVERRGMPSVGLADGTPVDMDHPYV FSDMTSYFTLLVGIYFPSVTGIMAGSNRSGDLRDAQKSIPTGTILAIATTSAVYISSV VLFGACIEGVVLRDKFGEAVNGNLVVGTLAWPSPWVIVIGSFFSTCGAGLQSLTGAPR LLQAISRDGIVPFLQVFGHGKANGEPTWALLLTACICEIGILIASLDEVAPILSMFFL MCYMFVNLACAVQTLLRTPNWRPRFRYYHWTLSFLGMSLCLALMFICSWYYALVAMLI AGLIYKYIEYRGAEKEWGDGIRGLSLSAARYALLRLEEGPPHTKNWRPQLLVLVRVDQ DQNVVHPQLLSLTSQLKAGKGLTIVGSVLEGTFLDNHPQAQRAEESIRRLMEAEKVKG FCQVVISSNLRDGVSHLIQSGGLGGLQHNTVLVGWPRNWRQKEDHQTWRNFIELVRET TAGHLALLVTKNVSMFPGNPERFSEGSIDVWWIVHDGGMLMLLPFLLRHHKVWRKCKM RIFTVAQMDDNSIQMKKDLTTFLYHLRITAEVEVVEMHESDISAYTYEKTLVMEQRSQ ILKQMHLTKNEREREIQSITDESRGSIRRKNPANTRLRLNVPEETACDNEEKPEEEVQ LIHDQSAPSCPSSSPSPGEEPEGEGETDPEKVHLTWTKDKSAAQKNKGPSPVSSEGIK DFFSMKPEWENLNQSNVRRMHTAVRLNEVIVNKSRDAKLVLLNMPGPPRNRNGDENYM EFLEVLTEQLDRVMLVRGGGREVITIYS

Rat KCC2 DNA (SEQ ID NO:5):

1 ccgctccacg gagagcaagc gacagagctc gagcaagcga gcgagcggcg aaggcgggca
61 gaggggcgcg ggcgaagagg cgcagccatc ccgagcccgg cgccgcgcag ccaccatgct
121 caacaacctg acggactgcg aggacggcga tgggggagcc aacccgggtg acggcaatcc
181 caaggagagc agccccttca tcaacagcac ggacacggag aaggggagag agtatgatgg
241 caggaacatg gccctgtttg aggaggagat ggacaccagc cccatggtat cctccctgct

301 cagtgg	getq qecaactac	a ccaacctee	- +		
361 aaacaa	gctg gccaactaca tgag ggcggaaag	a acaaccege	cagggaage	aaagagcaco	g aagaagcaga
		~ ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	. / 1/ 1/ 1: ATATA		
-			<i>(''</i> (')		
1621 ceteres	tgg gtcattgtca	taggctcttt	cttctctacc	tgcggagctq	gactacagag
1741 staste	ttt ggccatggca	aagccaacgg	agagccaacc	tgggcgctgc	tactaactac
		LUBILUI I I'AI	1 1 MATAA		
_		[4] [0] [7] [4] [4] [4]	(VIDAT		
			- ACCOT		
_ _			CICATEASA	+	
·			(12) (TCTTCCCC)		
			/12/17/2/2/2		
2001 Caagaac	gag cgggaacggg aat ccagccaaca	agatccagag	catcacagat	gaatctcggg	acticattica
3001 caacgag	gag aagccagaag	aggaggtgca	gctgatccat	qaccagagta	ctccaaata
orar agaggagg	egg aaccgtggtc	ccgggccggg	gagccccqaq	CCCatacata	aacaacaca aacaacaca
		_	5	J = 4000	acgcccc
				•	

```
3781 geogegetee ceeeggacee tggtegetga geoegggege egeteggetg egetatacat
 3841 agtgtacagg agacatcgag tgtattttta atgtccccat atttctgtaa actagaaacg
 3901 caacggacte ctcgccacgg ccgcgctctc cccgctgcgg gcgcccagga aggcggagac
 3961 ccgggaagcc agggttccct gcgctcccga gctgagagcc aagtgcttta aggccggcgc
 4021 teteetttee ettteetgte eaeggeeegg getteettet etteeteea gttettggeg
 4081 aacacaggtg aagcectgee eggtgeette gtggaggage aggegtetet eetetgttgg
 4141 cttgccgcct gctcccctg tcccgtggct cctcgccaaa gactgaattt gtggagctgg
 4201 agggcacacc ctccccactt tccttcctgg gacaggtgag gggccaatgc cagtctaggg
 4261 gccgactcac aggaggcctc gcgcagcctc ttggtcccca ctctgcaagt cctgcctggg
 4321 gacccagccc ccctggtggt tctggggcgg agctttgctg cctagcagca agtccttagt
 4381 tactgtctcc agataccagg acctggagta gggaatggag tcatatgggt tcagttgttc
 4441 ctggcgcttc tctgcccctt gctcccctc tccccctctc gtaggacaca aggactttgg
 4501 ctttcttaac tcatccttgg cgcttccgct ccaccacgcc cacctgtggg gaggagccct
 4561 cagccctaga gaggcgtttg gctggttccc ttcccccagg gcacgttact aagaggacag
 4621 gcactgcatg ctcctttaag cgccctctgg gactgggtac agtgcctcca gccccagggc
 4681 cctggtctgc gcacctagtt agacatcatt gcccactcca gggccagggc cactagctga
 4741 cctcaccacc tttttccttg agcccaaggc agagagagct gcagctggtg ccatctagac
 4801 aggetcaagt gtggccagtg gcagggctcg agggccactg ccctgttgct tggctcagga
 4861 cctctctgag atttgatggg gactggatat tcttccaggt agtagccatc aagtcggaag
 4921 tgttggaccc aggacctgac attccttcaa gactgccctc ccttgctgtg gttttgcctt
 4981 ttggggcaag agaggggctg ggcaaacggg gaggaggcag tatcaacacc gattagggaa
 5041 ccaaagttgc actacctggg cccagcctct ggttggcaag agcaaagttt ctgttgatga
 5101 aaacaaacag cccacaacaa caccccccc cccgttttct gtgctccatg tgcaatattt
 5161 gttatgaacc ttgtgtcgtt caagtcacct ttataatcac tgtagctaga tgttccatgt
 5221 ccatccaggt gactttactc tgagtgcaat atttcaatag cctggtagtg agaagagtgt
5341 ataggagagg ttgcaagcca aatcaagagt atttatcgtt attactatta ttattaggcc
 5401 tgcctttaat tttagtgttt cggtatttcg catcctgcct cggtattgat cgtgtgttct
 5461 ctgtgccaat atgcaaagga gaggatcagt tctttccttt actgttgaat gctcccattt
 5521 actgctttaa ggcttttact gtgttcattt tttagatacc tgtctg
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FIG. 11 (Continued)

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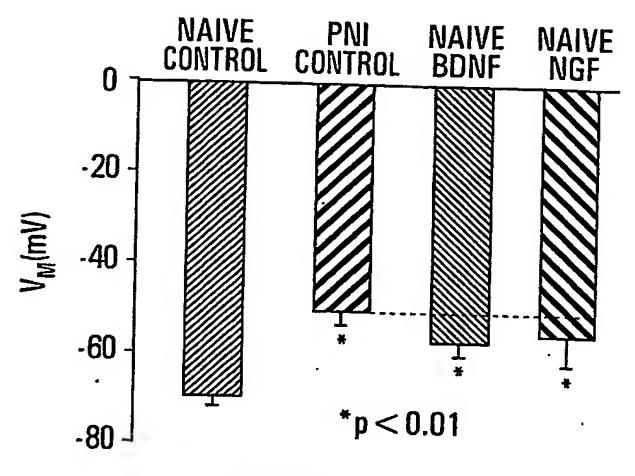


FIG. 12

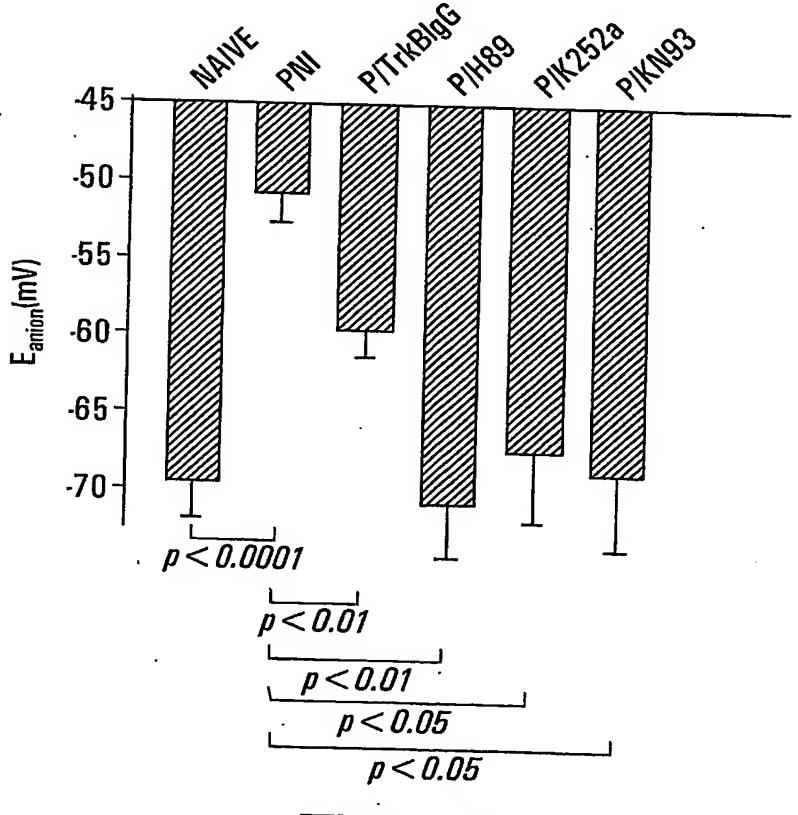
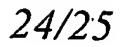


FIG. 13



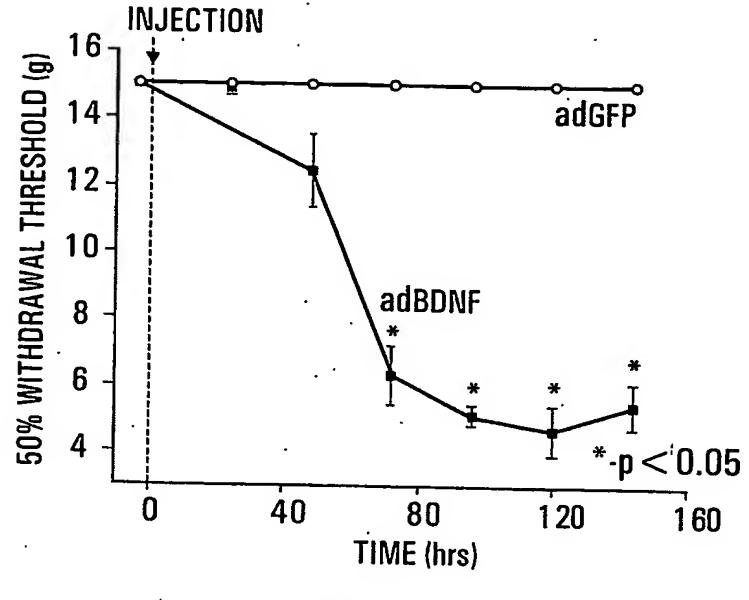


FIG. 14

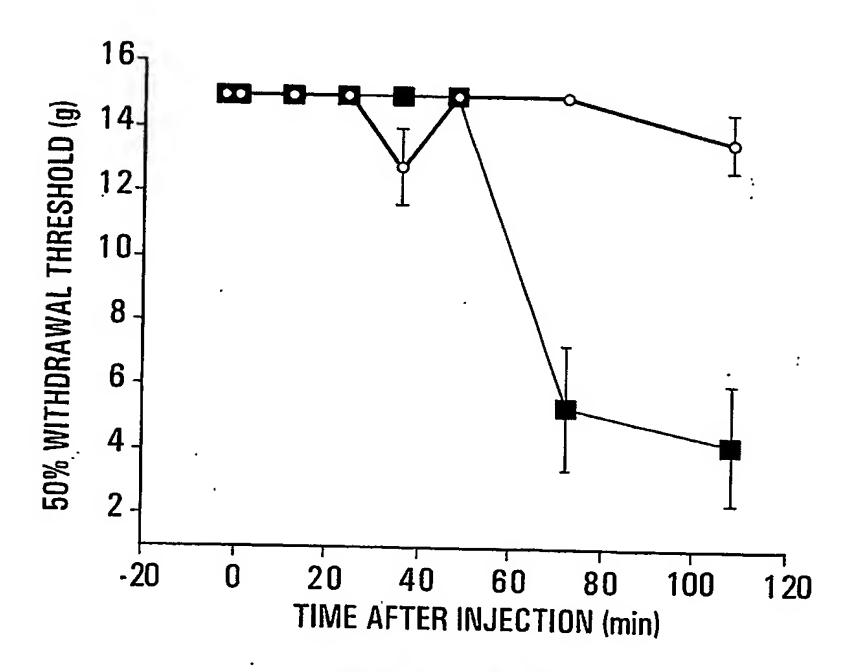
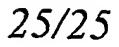
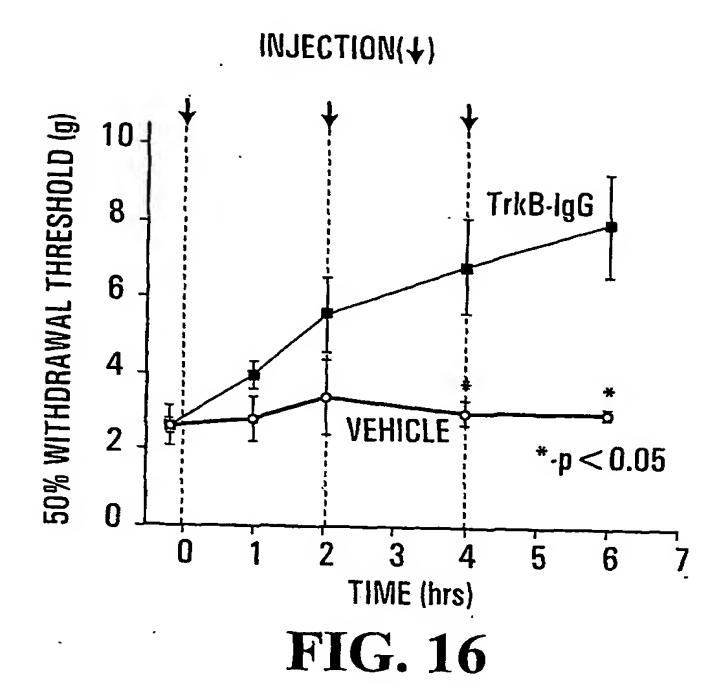


FIG. 15





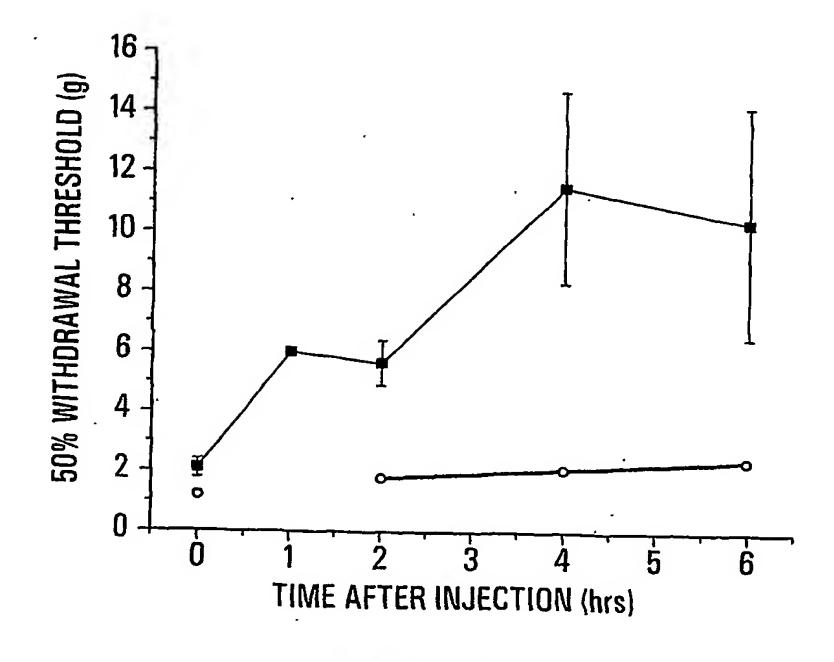


FIG. 17